

## **AMENDMENTS TO THE SPECIFICATION**

Please amend the specification as follows:

On page 1, immediately after the title insert:

### **RELATED APPLICATIONS**

This application is a US National Phase Application of PCT/ US2004/014540 filed on June 2, 2004, which claims priority to United States provisional patent application No. 60/475,402, filed June 2, 2003 and United States patent application No. 10/684,440, filed October 15, 2003, each of which is incorporated by reference in its entirety.

### **SEQUENCE LISTING**

A sequence listing is filed herewith in accordance with CFR 1.821 and is hereby incorporated by reference.

On page 10, line 29 to page 11, line 15:

Percent homology, sequence identity, or complementarity can be determined by, for example, the Gap program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, Madison WI), using default settings, which uses the algorithm of Smith and Waterman (Adv. Appl. Math., 1981, 2,482-489). In some embodiments, homology, sequence identity, or complementarity between the oligomeric compound and target is between about 50% to about 60%. In some embodiments, homology, sequence identity, or complementarity is between about 60% to about 70%. In other embodiments, homology, sequence identity, or complementarity is between about 70% and about 80%. In still other embodiments, homology, sequence identity, or complementarity is between about 80% and about 90%. In yet other embodiments, homology, sequence identity, or complementarity is about 90%, about 92%, about 94%, about 95%, about 96%, about 97%, about 98%, about 99%, or about 100%.